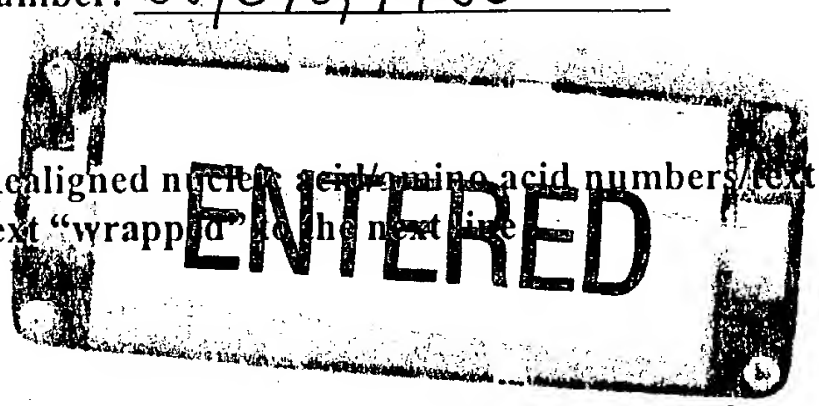
IFW16

CRF Errors Edited by the STIC Systems Branch

Serial Number: 08/390,740C

CRF Edit Date: 8/9/04

Edited by: [illegible]



___ Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line.

___ Corrected the SEQ ID NO. Sequence numbers edited were:

_____

___ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

_____

✓ Deleted: ___ invalid beginning/end-of-file text ; ___ page numbers

___ Inserted mandatory headings/numeric identifiers, specifically:

_____

___ Moved responses to same line as heading/numeric identifier, specifically:

_____

___ Other:

_____

_____

_____

Revised 09/09/2003

IFW16

RAW SEQUENCE LISTING DATE: 08/09/2004
PATENT APPLICATION: US/08/390,740C TIME: 15:12:52

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF4\08092004\H390740C.raw

SEQUENCE LISTING
C--> 5 (1) GENERAL INFORMATION:
7 (i) APPLICANT: Coleman, Roger
8 Bandman, Olga
9 Wilde, Craig G.
C--> 11 (ii) TITLE OF INVENTION: NEW CHEMOKINES EXPRESSED IN PANCREAS
13 (iii) NUMBER OF SEQUENCES: 11
15 (iv) CORRESPONDENCE ADDRESS:
16 (A) ADDRESSEE: Incyte Pharmaceuticals, Inc.
17 (B) STREET: 3174 Porter Drive
18 (C) CITY: Palo Alto
19 (D) STATE: CA
20 (E) COUNTRY: U.S.
21 (F) ZIP: 94304
23 (v) COMPUTER READABLE FORM:
24 (A) MEDIUM TYPE: Diskette
25 (B) COMPUTER: IBM Compatible
26 (C) OPERATING SYSTEM: DOS
27 (D) SOFTWARE: FastSEQ Version 1.5
29 (vi) CURRENT APPLICATION DATA:
C--> 30 (A) APPLICATION NUMBER: US/08/390,740C
C--> 31 (B) FILING DATE: 17-Feb-1995
33 (viii) ATTORNEY/AGENT INFORMATION:
34 (A) NAME: Luther, Barbara J.
35 (B) REGISTRATION NUMBER: 33,954
36 (C) REFERENCE/DOCKET NUMBER: PF-0027 US
38 (ix) TELECOMMUNICATION INFORMATION:
39 (A) TELEPHONE: 415-855-0555
40 (B) TELEFAX: 415-852-0195
43 (2) INFORMATION FOR SEQ ID NO: 1:
45 (i) SEQUENCE CHARACTERISTICS:
46 (A) LENGTH: 291 base pairs
47 (B) TYPE: nucleic acid
48 (C) STRANDEDNESS: single
49 (D) TOPOLOGY: linear
51 (ii) MOLECULE TYPE: cDNA
54 (vii) IMMEDIATE SOURCE:
55 (A) LIBRARY: Human Pancreas
56 (B) CLONE: 223187
58 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
60 ATGAAGGTCT CCGCAGCACT TCTGTGGCTG CTGCTCATAG CAGCTGCCTT CAGCCCCCAG 60
61 GGGCTCACTG GGCCAGCTTC TGTCCCAACC ACCTGCTGCT TTAACCTGGC CAATAGGAAG 120
62 ATACCCCTTC AGCGACTAGA GAGCTACAGG AGAATCACCA GTGGCAAATG TCCCCAGAAA 180

RAW SEQUENCE LISTING DATE: 08/09/2004
PATENT APPLICATION: **US/08/390,740C** TIME: 15:12:52

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\08092004\H390740C.raw**

GCTGTGATCT TCAAGACCAA ACTGGCCAAG GATATCTGTG CCGACCCCAA GAAGAAGTGG 240
GTGCAGGATT CCATGAAGTA TCTGGACCAA AAATCTCCAA CTCCAAAGCA 291

(2) INFORMATION FOR SEQ ID NO: 2:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 97 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(vii) IMMEDIATE SOURCE:
(A) LIBRARY: Human Pancreas
(B) CLONE: 223187

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

```
Met Lys Val Ser Ala Ala Leu Leu Trp Leu Leu Leu Ile Ala Ala Ala
 1               5                  10                  15
Phe Ser Pro Gln Gly Leu Thr Gly Pro Ala Ser Val Pro Thr Thr Cys
            20                  25                  30
Cys Phe Asn Leu Ala Asn Arg Lys Ile Pro Leu Gln Arg Leu Glu Ser
        35                  40                  45
Tyr Arg Arg Ile Thr Ser Gly Lys Cys Pro Gln Lys Ala Val Ile Phe
    50                  55                  60
Lys Thr Lys Leu Ala Lys Asp Ile Cys Ala Asp Pro Lys Lys Lys Trp
65                  70                  75                  80
Val Gln Asp Ser Met Lys Tyr Leu Asp Gln Lys Ser Pro Thr Pro Lys
                85                  90                  95
Pro
```

(2) INFORMATION FOR SEQ ID NO: 3:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 402 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(vii) IMMEDIATE SOURCE:
(A) LIBRARY: Human Pancreas
(B) CLONE: 226152

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

ATGGCTCAGT CACTGGCTCT GAGCCTCCTT ATCCTGGTTC TGGCCTTTGG CATCCCCAGG 60
ACCCAAGGCA GTGATGGAGG GGCTCAGGAC TGTTGCCTCA AGTACAGCCA AAGGAAGATT 120
CCCGCCAAGG TTGTCCGCAG CTACCGGAAG CAGGAACCAA GCTTAGGCTG CTCCATCCCA 180
GCTATCCTGT TCTTGCCCCG CAAGCGCTCT CAGGCAGAGC TATGTGCAGA CCCAAAGGAG 240
CTCTGGGTGC AGCAGCTGAT GCAGCATCTG GACAAGACAC CATCCCCACA GAAACCAGCC 300
CAGGGCTGCA GGAAGGACAG GGGGGCCTCC AAGACTGGCA AGAAAGGAAA GGGCTCCAAA 360
GGCTGCAAGA GGACTGAGCG GTCACAGACC CCTAAAGGGC CA 402

(2) INFORMATION FOR SEQ ID NO: 4:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 134 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single

RAW SEQUENCE LISTING DATE: 08/09/2004
PATENT APPLICATION: **US/08/390,740C** TIME: 15:12:52

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\08092004\H390740C.raw**

(D) TOPOLOGY: linear
(ii) MOLECULE TYPE: peptide
(vii) IMMEDIATE SOURCE:
(A) LIBRARY: Human Pancreas
(B) CLONE: 226152
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

```
Met Ala Gln Ser Leu Ala Leu Ser Leu Leu Ile Leu Val Leu Ala Phe
 1               5                  10                  15
Gly Ile Pro Arg Thr Gln Gly Ser Asp Gly Gly Ala Gln Asp Cys Cys
            20                  25                  30
Leu Lys Tyr Ser Gln Arg Lys Ile Pro Ala Lys Val Val Arg Ser Tyr
        35                  40                  45
Arg Lys Gln Glu Pro Ser Leu Gly Cys Ser Ile Pro Ala Ile Leu Phe
    50                  55                  60
Leu Pro Arg Lys Arg Ser Gln Ala Glu Leu Cys Ala Asp Pro Lys Glu
65                  70                  75                  80
Leu Trp Val Gln Gln Leu Met Gln His Leu Asp Lys Thr Pro Ser Pro
                85                  90                  95
Gln Lys Pro Ala Gln Gly Cys Arg Lys Asp Arg Gly Ala Ser Lys Thr
            100                 105                 110
Gly Lys Lys Gly Lys Gly Ser Lys Gly Cys Lys Arg Thr Glu Arg Ser
        115                 120                 125
Gln Thr Pro Lys Gly Pro
    130
```

(2) INFORMATION FOR SEQ ID NO: 5:
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 97 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
(ii) MOLECULE TYPE: peptide
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

```
Met Lys Val Ser Ala Ala Leu Leu Ala Leu Leu Leu Ile Ala Ala Ala
 1               5                  10                  15
Phe Cys Pro Gln Gly Leu Ala Gln Pro Asp Gly Val Asp Thr Pro Thr
            20                  25                  30
Thr Cys Cys Phe Asn Tyr Ile Asn Arg Lys Ile Pro Arg Gln Arg Leu
        35                  40                  45
Glu Ser Tyr Arg Arg Ile Thr Ser Ser Lys Cys Ser Lys Pro Ala Val
    50                  55                  60
Ile Phe Lys Thr Lys Arg Ala Lys Gln Val Cys Ala Asp Pro Lys Glu
65                  70                  75                  80
Lys Trp Val Gln Asp Ser Met Lys His Leu Asp Lys Gln Thr Pro Lys
                85                  90                  95
Pro
```

(2) INFORMATION FOR SEQ ID NO: 6:
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 92 amino acids
(B) TYPE: amino acid

RAW SEQUENCE LISTING DATE: 08/09/2004
PATENT APPLICATION: **US/08/390,740C** TIME: 15:12:52

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\08092004\H390740C.raw**

(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
(ii) MOLECULE TYPE: peptide
(vii) IMMEDIATE SOURCE:
(A) LIBRARY: GenBank
(B) CLONE: MIP-1a
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:

Met Gln Val Ser Thr Ala Ala Leu Ala Val Leu Leu Cys Thr Met Ala
1 5 10 15
Leu Cys Asn Gln Phe Ser Ala Ser Leu Ala Ala Asp Thr Pro Thr Ala
20 25 30
Cys Cys Phe Ser Tyr Thr Ser Arg Gln Ile Pro Gln Asn Phe Ile Ala
35 40 45
Asp Tyr Phe Glu Thr Ser Ser Gln Cys Ser Lys Pro Gly Val Ile Phe
50 55 60
Leu Thr Lys Arg Ser Arg Gln Val Cys Ala Asp Pro Ser Glu Glu Trp
65 70 75 80
Val Gln Lys Tyr Val Ser Asp Leu Glu Leu Ser Ala
85 90

(2) INFORMATION FOR SEQ ID NO: 7:
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 92 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
(ii) MOLECULE TYPE: peptide
(vii) IMMEDIATE SOURCE:
(A) LIBRARY: GenBank
(B) CLONE: MIP-1b
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

Met Lys Leu Cys Val Thr Val Leu Ser Leu Leu Met Leu Val Ala Ala
1 5 10 15
Phe Cys Ser Pro Ala Leu Ser Ala Pro Met Gly Ser Asp Pro Pro Thr
20 25 30
Ala Cys Cys Phe Ser Tyr Thr Ala Arg Lys Leu Pro Arg Asn Phe Val
35 40 45
Val Asp Tyr Tyr Glu Thr Ser Ser Leu Cys Ser Gln Pro Ala Val Val
50 55 60
Phe Gln Thr Lys Arg Ser Lys Gln Val Cys Ala Asp Pro Ser Glu Ser
65 70 75 80
Trp Val Gln Glu Tyr Val Tyr Asp Leu Glu Leu Asn
85 90

(2) INFORMATION FOR SEQ ID NO: 8:
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 91 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
(ii) MOLECULE TYPE: peptide

RAW SEQUENCE LISTING DATE: 08/09/2004
PATENT APPLICATION: **US/08/390,740C** TIME: 15:12:52

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\08092004\H390740C.raw**

(vii) IMMEDIATE SOURCE:
(A) LIBRARY: GenBank
(B) CLONE: RANTES

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

Met Lys Val Ser Ala Ala Arg Leu Ala Val Ile Leu Ile Ala Thr Ala
1 5 10 15
Leu Cys Ala Pro Ala Ser Ala Ser Pro Tyr Ser Ser Asp Thr Thr Pro
20 25 30
Cys Cys Phe Ala Tyr Ile Ala Arg Pro Leu Pro Arg Ala His Ile Lys
35 40 45
Glu Tyr Phe Tyr Thr Ser Gly Lys Cys Ser Asn Pro Ala Val Val Phe
50 55 60
Val Thr Arg Lys Asn Arg Gln Val Cys Ala Asn Pro Glu Lys Lys Trp
65 70 75 80
Val Arg Glu Tyr Ile Asn Ser Leu Glu Met Ser
85 90

(2) INFORMATION FOR SEQ ID NO: 9:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 99 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(vii) IMMEDIATE SOURCE:
(A) LIBRARY: GenBank
(B) CLONE: MCP-1

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:

Met Lys Val Ser Ala Ala Leu Leu Cys Leu Leu Leu Ile Ala Ala Thr
1 5 10 15
Phe Ile Pro Gln Gly Leu Ala Gln Pro Asp Ala Ile Asn Ala Pro Val
20 25 30
Thr Cys Cys Tyr Asn Phe Thr Asn Arg Lys Ile Ser Val Gln Arg Leu
35 40 45
Ala Ser Tyr Arg Arg Ile Thr Ser Ser Lys Cys Pro Lys Glu Ala Val
50 55 60
Ile Phe Lys Thr Ile Val Ala Lys Glu Ile Cys Ala Asp Pro Lys Gln
65 70 75 80
Lys Trp Val Gln Asp Ser Met Asp His Leu Asp Lys Gln Thr Gln Thr
85 90 95
Pro Lys Thr

(2) INFORMATION FOR SEQ ID NO: 10:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 77 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(vii) IMMEDIATE SOURCE:
(A) LIBRARY: GenBank

VERIFICATION SUMMARY DATE: 08/09/2004
PATENT APPLICATION: **US/08/390,740C** TIME: 15:12:53

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\08092004\H390740C.raw**

M:220 C: Keyword misspelled or invalid format, [(1) GENERAL INFORMATION:]
1 M:220 C: Keyword misspelled or invalid format, [(ii) TITLE OF INVENTION:]
0 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
1 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
54 M:238 W: Alpha Fields not Ordered, Reordered [(vi) ORIGINAL SOURCE:] of (2)

IFW16

RAW SEQUENCE LISTING DATE: 08/06/2004
PATENT APPLICATION: US/08/390,740C TIME: 14:29:21

Input Set : A:\Sequence Listings.txt
Output Set: N:\CRF4\08062004\H390740C.raw

Does Not Comply
Corrected Diskette Needed

SEQUENCE LISTING

W--> 1 PF-0027 US SUBSTITUTE SHEET
C--> 5 (1) GENERAL INFORMATION:
7 (i) APPLICANT: Coleman, Roger
8 Bandman, Olga
9 Wilde, Craig G.
C--> 11 (ii) TITLE OF INVENTION: NEW CHEMOKINES EXPRESSED IN PANCREAS
13 (iii) NUMBER OF SEQUENCES: 11
15 (iv) CORRESPONDENCE ADDRESS:
16 (A) ADDRESSEE: Incyte Pharmaceuticals, Inc.
17 (B) STREET: 3174 Porter Drive
18 (C) CITY: Palo Alto
19 (D) STATE: CA
20 (E) COUNTRY: U.S.
21 (F) ZIP: 94304
23 (v) COMPUTER READABLE FORM:
24 (A) MEDIUM TYPE: Diskette
25 (B) COMPUTER: IBM Compatible
26 (C) OPERATING SYSTEM: DOS
27 (D) SOFTWARE: FastSEQ Version 1.5
29 (vi) CURRENT APPLICATION DATA:
C--> 30 (A) APPLICATION NUMBER: US/08/390,740C
C--> 31 (B) FILING DATE: 17-Feb-1995
33 (viii) ATTORNEY/AGENT INFORMATION:
34 (A) NAME: Luther, Barbara J.
35 (B) REGISTRATION NUMBER: 33,954
36 (C) REFERENCE/DOCKET NUMBER: PF-0027 US
38 (ix) TELECOMMUNICATION INFORMATION:
39 (A) TELEPHONE: 415-855-0555
40 (B) TELEFAX: 415-852-0195
43 (2) INFORMATION FOR SEQ ID NO: 1:
45 (i) SEQUENCE CHARACTERISTICS:
46 (A) LENGTH: 291 base pairs
47 (B) TYPE: nucleic acid
48 (C) STRANDEDNESS: single
49 (D) TOPOLOGY: linear
51 (ii) MOLECULE TYPE: cDNA
54 (vii) IMMEDIATE SOURCE:
55 (A) LIBRARY: Human Pancreas
56 (B) CLONE: 223187
58 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
60 ATGAAGGTCT CCGCAGCACT TCTGTGGCTG CTGCTCATAG CAGCTGCCTT CAGCCCCCAG 60
61 GGGCTCACTG GGCCAGCTTC TGTCCCAACC ACCTGCTGCT TTAACCTGGC CAATAGGAAG 120

RAW SEQUENCE LISTING DATE: 08/06/2004
PATENT APPLICATION: **US/08/390,740C** TIME: 14:29:21

Input Set : **A:\Sequence Listings.txt**
Output Set: **N:\CRF4\08062004\H390740C.raw**

ATACCCCTTC AGCGACTAGA GAGCTACAGG AGAATCACCA GTGGCAAATG TCCCCAGAAA 180
GCTGTGATCT TCAAGACCAA ACTGGCCAAG GATATCTGTG CCGACCCCAA GAAGAAGTGG 240
GTGCAGGATT CCATGAAGTA TCTGGACCAA AAATCTCCAA CTCCAAAGCC A 291

(2) INFORMATION FOR SEQ ID NO: 2:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 97 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(vii) IMMEDIATE SOURCE:
(A) LIBRARY: Human Pancreas
(B) CLONE: 223187

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

```
Met Lys Val Ser Ala Ala Leu Leu Trp Leu Leu Leu Ile Ala Ala Ala
 1               5                  10                  15
Phe Ser Pro Gln Gly Leu Thr Gly Pro Ala Ser Val Pro Thr Thr Cys
            20                  25                  30
Cys Phe Asn Leu Ala Asn Arg Lys Ile Pro Leu Gln Arg Leu Glu Ser
        35                  40                  45
Tyr Arg Arg Ile Thr Ser Gly Lys Cys Pro Gln Lys Ala Val Ile Phe
    50                  55                  60
Lys Thr Lys Leu Ala Lys Asp Ile Cys Ala Asp Pro Lys Lys Lys Trp
65                  70                  75                  80
Val Gln Asp Ser Met Lys Tyr Leu Asp Gln Lys Ser Pro Thr Pro Lys
                85                  90                  95
Pro
```

(2) INFORMATION FOR SEQ ID NO: 3:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 402 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: cDNA

(vii) IMMEDIATE SOURCE:
(A) LIBRARY: Human Pancreas
(B) CLONE: 226152

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

ATGGCTCAGT CACTGGCTCT GAGCCTCCTT ATCCTGGTTC TGGCCTTTGG CATCCCCAGG 60
ACCCAAGGCA GTGATGGAGG GGCTCAGGAC TGTTGCCTCA AGTACAGCCA AAGGAAGATT 120
CCCGCCAAGG TTGTCCGCAG CTACCGGAAG CAGGAACCAA GCTTAGGCTG CTCCATCCCA 180
GCTATCCTGT TCTTGCCCCG CAAGCGCTCT CAGGCAGAGC TATGTGCAGA CCCAAAGGAG 240
CTCTGGGTGC AGCAGCTGAT GCAGCATCTG GACAAGACAC CATCCCCACA GAAACCAGCC 300
CAGGGCTGCA GGAAGGACAG GGGGGCCTCC AAGACTGGCA AGAAAGGAAA GGGCTCCAAA 360
GGCTGCAAGA GGACTGAGCG GTCACAGACC CCTAAAGGGC CA 402

(2) INFORMATION FOR SEQ ID NO: 4:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 134 amino acids
(B) TYPE: amino acid

(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
(ii) MOLECULE TYPE: peptide
(vii) IMMEDIATE SOURCE:
(A) LIBRARY: Human Pancreas
(B) CLONE: 226152
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

```
Met Ala Gln Ser Leu Ala Leu Ser Leu Leu Ile Leu Val Leu Ala Phe
 1               5                  10                  15
Gly Ile Pro Arg Thr Gln Gly Ser Asp Gly Gly Ala Gln Asp Cys Cys
            20                  25                  30
Leu Lys Tyr Ser Gln Arg Lys Ile Pro Ala Lys Val Val Arg Ser Tyr
        35                  40                  45
Arg Lys Gln Glu Pro Ser Leu Gly Cys Ser Ile Pro Ala Ile Leu Phe
    50                  55                  60
Leu Pro Arg Lys Arg Ser Gln Ala Glu Leu Cys Ala Asp Pro Lys Glu
65                  70                  75                  80
Leu Trp Val Gln Gln Leu Met Gln His Leu Asp Lys Thr Pro Ser Pro
                85                  90                  95
Gln Lys Pro Ala Gln Gly Cys Arg Lys Asp Arg Gly Ala Ser Lys Thr
            100                 105                 110
Gly Lys Lys Gly Lys Gly Ser Lys Gly Cys Lys Arg Thr Glu Arg Ser
        115                 120                 125
Gln Thr Pro Lys Gly Pro
    130
```

(2) INFORMATION FOR SEQ ID NO: 5:
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 97 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
(ii) MOLECULE TYPE: peptide
(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

```
Met Lys Val Ser Ala Ala Leu Leu Ala Leu Leu Leu Ile Ala Ala Ala
 1               5                  10                  15
Phe Cys Pro Gln Gly Leu Ala Gln Pro Asp Gly Val Asp Thr Pro Thr
            20                  25                  30
Thr Cys Cys Phe Asn Tyr Ile Asn Arg Lys Ile Pro Arg Gln Arg Leu
        35                  40                  45
Glu Ser Tyr Arg Arg Ile Thr Ser Ser Lys Cys Ser Lys Pro Ala Val
    50                  55                  60
Ile Phe Lys Thr Lys Arg Ala Lys Gln Val Cys Ala Asp Pro Lys Glu
65                  70                  75                  80
Lys Trp Val Gln Asp Ser Met Lys His Leu Asp Lys Gln Thr Pro Lys
                85                  90                  95
Pro
```

(2) INFORMATION FOR SEQ ID NO: 6:
(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 92 amino acids

RAW SEQUENCE LISTING DATE: 08/06/2004
PATENT APPLICATION: **US/08/390,740C** TIME: 14:29:21

Input Set : **A:\Sequence Listings.txt**
Output Set: **N:\CRF4\08062004\H390740C.raw**

(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(vii) IMMEDIATE SOURCE:
(A) LIBRARY: GenBank
(B) CLONE: MIP-1a

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:

```
Met Gln Val Ser Thr Ala Ala Leu Ala Val Leu Leu Cys Thr Met Ala
 1               5                  10                  15
Leu Cys Asn Gln Phe Ser Ala Ser Leu Ala Ala Asp Thr Pro Thr Ala
            20                  25                  30
Cys Cys Phe Ser Tyr Thr Ser Arg Gln Ile Pro Gln Asn Phe Ile Ala
        35                  40                  45
Asp Tyr Phe Glu Thr Ser Ser Gln Cys Ser Lys Pro Gly Val Ile Phe
    50                  55                  60
Leu Thr Lys Arg Ser Arg Gln Val Cys Ala Asp Pro Ser Glu Glu Trp
65                  70                  75                  80
Val Gln Lys Tyr Val Ser Asp Leu Glu Leu Ser Ala
                85                  90
```

(2) INFORMATION FOR SEQ ID NO: 7:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 92 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(vii) IMMEDIATE SOURCE:
(A) LIBRARY: GenBank
(B) CLONE: MIP-1b

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

```
Met Lys Leu Cys Val Thr Val Leu Ser Leu Leu Met Leu Val Ala Ala
 1               5                  10                  15
Phe Cys Ser Pro Ala Leu Ser Ala Pro Met Gly Ser Asp Pro Pro Thr
            20                  25                  30
Ala Cys Cys Phe Ser Tyr Thr Ala Arg Lys Leu Pro Arg Asn Phe Val
        35                  40                  45
Val Asp Tyr Tyr Glu Thr Ser Ser Leu Cys Ser Gln Pro Ala Val Val
    50                  55                  60
Phe Gln Thr Lys Arg Ser Lys Gln Val Cys Ala Asp Pro Ser Glu Ser
65                  70                  75                  80
Trp Val Gln Glu Tyr Val Tyr Asp Leu Glu Leu Asn
                85                  90
```

(2) INFORMATION FOR SEQ ID NO: 8:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 91 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

RAW SEQUENCE LISTING DATE: 08/06/2004
PATENT APPLICATION: **US/08/390,740C** TIME: 14:29:21

Input Set : **A:\Sequence Listings.txt**
Output Set: **N:\CRF4\08062004\H390740C.raw**

(ii) MOLECULE TYPE: peptide

(vii) IMMEDIATE SOURCE:
(A) LIBRARY: GenBank
(B) CLONE: RANTES

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

```
Met Lys Val Ser Ala Ala Arg Leu Ala Val Ile Leu Ile Ala Thr Ala
 1               5                  10                  15
Leu Cys Ala Pro Ala Ser Ala Ser Pro Tyr Ser Ser Asp Thr Thr Pro
            20                  25                  30
Cys Cys Phe Ala Tyr Ile Ala Arg Pro Leu Pro Arg Ala His Ile Lys
        35                  40                  45
Glu Tyr Phe Tyr Thr Ser Gly Lys Cys Ser Asn Pro Ala Val Val Phe
    50                  55                  60
Val Thr Arg Lys Asn Arg Gln Val Cys Ala Asn Pro Glu Lys Lys Trp
65                  70                  75                  80
Val Arg Glu Tyr Ile Asn Ser Leu Glu Met Ser
                85                  90
```

(2) INFORMATION FOR SEQ ID NO: 9:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 99 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(vii) IMMEDIATE SOURCE:
(A) LIBRARY: GenBank
(B) CLONE: MCP-1

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:

```
Met Lys Val Ser Ala Ala Leu Leu Cys Leu Leu Leu Ile Ala Ala Thr
 1               5                  10                  15
Phe Ile Pro Gln Gly Leu Ala Gln Pro Asp Ala Ile Asn Ala Pro Val
            20                  25                  30
Thr Cys Cys Tyr Asn Phe Thr Asn Arg Lys Ile Ser Val Gln Arg Leu
        35                  40                  45
Ala Ser Tyr Arg Arg Ile Thr Ser Ser Lys Cys Pro Lys Glu Ala Val
    50                  55                  60
Ile Phe Lys Thr Ile Val Ala Lys Glu Ile Cys Ala Asp Pro Lys Gln
65                  70                  75                  80
Lys Trp Val Gln Asp Ser Met Asp His Leu Asp Lys Gln Thr Gln Thr
                85                  90                  95
Pro Lys Thr
```

(2) INFORMATION FOR SEQ ID NO: 10:

(i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 77 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(vii) IMMEDIATE SOURCE:

VERIFICATION SUMMARY DATE: 08/06/2004
PATENT APPLICATION: **US/08/390,740C** TIME: 14:29:22

Input Set : **A:\Sequence Listings.txt**
Output Set: **N:\CRF4\08062004\H390740C.raw**

L:1 M:244 W: Invalid beginning of sequence listing, Line=[PF-0027 US SUBSTITUTE SHEET],
General Header Line Not Processed!
L:5 M:220 C: Keyword misspelled or invalid format, [(1) GENERAL INFORMATION:]
L:11 M:220 C: Keyword misspelled or invalid format, [(ii) TITLE OF INVENTION:]
L:30 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:31 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:354 M:238 W: Alpha Fields not Ordered, Reordered [(vi) ORIGINAL SOURCE:] of (2)